

RAW SEQUENCE LISTING ERROR REPORT

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NOTICE TO COMPLY FOR CRESUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2:16 mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax) PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TOREDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.1. PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

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plicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom. Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- EES:Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>, EFS Submission User Manual e e PAVB)
- 2. aU.S. Postal Service: U.S. Patent and Trademark Office; Box Sequence, P.O. Box 2327, Arlington, VA 22202

3. Hand Carry directly to:

- U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
 - U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
 - 4. Federal Express, United Parcel Service, r other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 01/29/2002



OIPE

DATE: 02/19/2002 RAW SEQUENCE LISTING TIME: 14:36:23 PATENT APPLICATION: US/09/928,847A Input Set : A:\10022204.ST25.txt Output Set: N:\CRF3\02192002\I928847A.raw 3 <110> APPLICANT: Novozymes A/S Jorgensen, Steen T 5 Rasmussen, Michael D Andersen, Jens Tonne Olsen, Carsten 9 <120> TITLE OF INVENTION: Multiple Insertion of Genes Doss Not Comply 11 <130> FILE REFERENCE: 10022.204-US Corrected Diskette Needec 13 <140> CURRENT APPLICATION NUMBER: 09/928847A 14 <141> CURRENT FILING DATE: 2001-08-13 16 <160> NUMBER OF SEQ ID NOS: 50 18 <170> SOFTWARE: PatentIn version 3.1 20 <210> SEQ ID NO: 1 21 <211> LENGTH: 30 22 <212> TYPE: DNA 23 <213> ORGANISM: Artificial Sequence 25 <220> FEATURE: 26 <223> OTHER INFORMATION: Primer 28 <400> SEQUENCE: 1 29 gactaagctt ctgcatagtg agagaagacg 30 32 <210> SEQ ID NO: 2 33 <211> LENGTH: 67 34 <212> TYPE: DNA 35 <213> ORGANISM: Artificial Sequence 37 <220> FEATURE: 38 <223> OTHER INFORMATION: Primer 40 <400> SEQUENCE: 2 41 gactgaattc agatctgcgg ccgcacgcgt gtcgacagta ctgaaataga ggaaaaaata 60 43 agttttc 67 46 <210> SEQ ID NO: 3 47 <211> LENGTH: 33 48 <212> TYPE: DNA 49 <213> ORGANISM: Artificial Sequence 51 <220> FEATURE: 52 <223> OTHER INFORMATION: Primer 54 <400> SEQUENCE: 3 33 55 gactgaattc cgtatccatt cctgcgatat gag 58 <210> SEQ ID NO: 4 59 <211> LENGTH: 41 60 <212> TYPE: DNA 61 <213> ORGANISM: Artificial Sequence 63 <220> FEATURE: 64 <223> OTHER INFORMATION: Primer

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176 gctttctgcc tgcggccaaa aatcgcaaga agatgttgtg acggggctcg acaagaaggc
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178 aaaagaatac acgtcctata aggcaaaagc gaaaatgacc attgaaacgg ggaatgaccc
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                                                                           300
180 gcaggagtac aacgtggaaa tctggcataa aaaaccttct ctttaccggg tctatttgga
182 aaacccqaaa aaaqaccaqa qccaqqtqat cttgcgcaat gaaaacggcg tgtttgtttt
                                                                           360
184 gactccgtcg ctgaataaaa gcttccgctt tcacagcgac tggcccaata acagcagcca
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                                                                           540
188 cqcaaaqqat tcaaaataca tttttqaaac qaaaacgaat tatcagcata atcagatgct
190 gccgactcag gaaatcgttt tccataaaaa gaccatggct ccttcatcgg ttaaagtgat
                                                                           600
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198 cqtqaaaaaq cttqaaqaqa aaqaqatqqc qactqaagac ggcaaacgga tcgtcatcac
                                                                           840
                                                                           900
200 atatggcggt gaaaaatcct ttacattgat tcaggaaaaa gcccgcgtcg ccaaaacatc
202 cactteegta tecatgaacg gagageeegt tgacetegge tteaeggteg gegeactgae
                                                                          960
204 ggataaatcg ttgtcatgga catatgacgg agtcgattac tttatctcat cagaagatct
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206 ttctcaagat gaacttctga tggttgcaaa aagcatgcag ggacagtctt cgaaatagac
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208 tgtgccgtat ccggcagcct gttttccgcc cggaagcgga aagcaggctt ttttatattt
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210 gcgtcgcaag cgtatgattt cgacagcttt tccgtaaaat gtataccgtg ccagcaattt
                                                                         1200
212 ttettttgtt cagggetgat gatecegtge aaaattteee ttteteegaa etttttagta
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214 tgatgggaag gacgagtgaa acaaggaaca ggaagtgtca tg atg agc tta aaa
                                                                         1314
                                                    Met Ser Leu Lys
215
216
                                                                         1362
218 cca ttc tat aga aag aca tgg gcc gaa atc gat tta acg gct tta aaa
219 Pro Phe Tyr Arg Lys Thr Trp Ala Glu Ile Asp Leu Thr Ala Leu Lys
                                             15
                        10
                                                                         1410
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RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/928,847A
DATE: 02/19/2002
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Input Set : A:\10022204.ST25.txt

Output Set: N:\CRF3\02192002\1928847A.raw

222	Glu	Aan	บรา	λκα	λan	Mot	Luc	λνα	Uic	Tla	Glv	Glu	Hic	Va 1	Δrσ	T.611	
223	GIU	ASII	vai	AIG	25	Mec	цуѕ	AIG	птэ	30	GIY	Gru	1113	Vai	35	пеа	
	atg	qcc	gtc	gtt		gcg	aat	gcc	tac	gga	cac	ggg	gat	gca	cag	gta	1458
	Met																
228				40					45					50			
	gcg																1506
231	Ala	Lys	Ala	Ala	Leu	Ala	Glu	Gly	Ala	Ser	Ile	Leu		Val	Ala	Leu	
232		*	55					60					65				4554
	ttg																1554
	Leu		Glu	Ala	Leu	Ser		Arg	Ата	GLn	GTĀ		Glu	GLU	Pro	ire	
236	ctt	70	a+ a	~~+	~~~	~+ ~	75	200	<i>a</i>	+ > +	~~~	80	2++	aaa	aaa	<i>a</i> 2 2	1602
	Leu																1002
240		vai	Leu	СТУ	Ата	90	FIU	1111	Giu	- y -	95	Ser	110	ALU	niu.	100	
	aag	cac	att	atc	ata		aac	tac	tida	atc		t.aa	ct.a	aaa	gac		1650
	Lys																
244		5			105		1	-1-		110	2	_		1	115		
	ctc	ggt	ttt	ctq	aat	gag	qcc	gaa	gct	cct	ctt	gaa	tat	cat	ttg	aag	1698
	Leu																
248		-		120					125					130			
	atc																1746
	Ile	Asp		Gly	Met	Gly	Arg		Gly	Cys	Lys	Thr		Glu	Glu	Ile	
252			135					140					145				4504
																acg	1794
	Lys		Met	Met	GIU	Met		GIU	ser	Asn	Asp	Lуs 160	Leu	Asn	Cys	Thr	
256	~~~	150	++0	a a t	ast	++0	155	200	aca	a a a	a a a		asa	200	rat	tat	1842
	ggc Gly																1042
	165	Val	FIIC	1111	1113	170	AIG	1111	AIG	LSP	175	цуз	1150	1111	nsp	180	
	ttc	aac	at.σ	cat	ctt		cac	ttt	aaa	σασ		atc	agc	ccc	ttc		1890
	Phe																
264					185	-	_		-	190					195		
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267	Leu	Asp	Arg	Leu	Met	Val	His	Ser	Ser	Asn	Ser	Ala	Ala	Gly	Leu	Arg	
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	ttc																1986
	Phe	Arg		Gln	Leu	Phe	Asn		Val	Arg	Phe	GLY		GTA	Met	Tyr	
272			215					220					225				2034
	ggt Gly																2034
275	СТУ	230	Ата	PIO	ser	1 111	235	тте	гуѕ	ASP	GIU	240	PIO	PHE	ALY	nea	
	cgg	-	ata	+++	tca	ctt		acc	паа	ctc	acc		ata	aaa	aaa	att	2082
	Arg																2002
	245					250	,_ J. D				255		·	-1-	-1-	260	
	aaa	aaa	qqc	gag	agc		agc	tac	qqq	gcg		tat	aca	gct	cag	cgc	2130
	Lys																
284		-	-		265			_	_	270		_			275	-	
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287	Asp	Glu	${\tt Trp}$	Ile	Gly	Thr	Val	Pro	Val	Gly	Tyr	Ala	Asp	Gly	Trp	Leu	

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		output oct	. H. (CRI 3	(02131	2002 (13.		- ••		
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	Arg Arg Leu								
292	295		300		-	305	<i>y</i> .	4	
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	Ile Ala Gly								
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	Glu Tyr Pro								
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	Glu Trp Ile								
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306	tat gaa att	acc tgt at	g ata agt	tca a	agg gtg	ccc cgt	atg ttt	ttg	2418
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308		360		365			370		
	gaa aat ggg								2466
311	Glu Asn Gly	Ser Ile Me	t Glu Ile	Arg A	Asn Pro	Ile Leu	Pro Asp	Gln	
312	375		380			385			
314	tcc tgaaaat	tga tgaatta	gcg gaaaa	acaac	tttgctt	gcg aaaa	agaataa		2519
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	ttgatgcgaa cgattgacaa								3179
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	tgctcgaggc								3299
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	<211> LENGT								
	<211> EERG1:				•				
	<213> ORGAN		ıs lichen	iformi	is			- 1	
	<220> FEATU		1011011						`) .
	<221> NAME/I		eature					In	, is an
	<222> LOCAT							. 00	nero sel
355	<223> OTHER	INFORMATIO	v: n deno	tes an	n undete	ermined n	ucleotid	le)	
	<400> SEQUE								
	Met Ser Leu		e Tyr Arg	Lys I	Thr Trp	Ala Glu	Ile Asp	Leu /	Elquerce
360		5			10		15	Ĺ	lease correct is type of liver
	Thr Ala Leu	Lys Glu Ası	n Val Arg	Asn M	Met Lys	Arg His	Ile Gly	Glu	1 t
364		20		25			30		esse concer
367	His Val Arg	Leu Met Ala	a Val Val	Lys A	Ala Asn	Ala Tyr	Gly His	Gly	+ true of orner
								T.W.	2 July 1

FJI)

Use of n and/or Xaa has been detected in the Geouence Listing. Review the Sequence Listing to insure a corresponding explanation is presented in the <220> to <223> fields of each sequence using n or Xaa.

VERIFICATION SUMMARY

DATE: 02/19/2002 TIME: 14:36:24

PATENT APPLICATION: US/09/928,847A

Input Set : $A:\10022204.ST25.txt$

Output Set: N:\CRF3\02192002\I928847A.raw

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